

## SEQUENCE LISTING

<110> Glimcher, Laurie H. Szabo, Susanne, J.

<120> T-BET COMPOSITIONS AND METHODS OF USE THEREOF <130> HUI-040CP <140> US 10/008264 <141> 2001-12-03 <150> PCT/US00/15345 <151> 2000-06-01 <150> US 60/137085 <151> 1999-06-02 <160> 4 <170> PatentIn Ver. 2.0 <210> 1 <211> 1608 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1605) <400> 1 atg ggc atc gtg gag ccg ggt tgc gga gac atg ctg acg ggc acc gag Met Gly Ile Val Glu Pro Gly Cys Gly Asp Met Leu Thr Gly Thr Glu ccg atg ccg ggg agc gac gag ggc cgg gcg cct ggc gcc gac ccg cag Pro Met Pro Gly Ser Asp Glu Gly Arg Ala Pro Gly Ala Asp Pro Gln cae ege tae tte tae eeg gag eeg gge geg eag gae geg gae gag egt 144 His Arg Tyr Phe Tyr Pro Glu Pro Gly Ala Gln Asp Ala Asp Glu Arg cgc ggg ggc ggc agc ctg ggg tct ccc tac ccg ggg ggc gcc ttg gtg 192 Arg Gly Gly Ser Leu Gly Ser Pro Tyr Pro Gly Gly Ala Leu Val ccc gcc ccg ccg agc cgc ttc ctt gga gcc tac gcc tac ccg ccg cga 240 Pro Ala Pro Pro Ser Arg Phe Leu Gly Ala Tyr Ala Tyr Pro Pro Arg 65 ccc cag gcg gcc ggc ttc ccc ggc gcg ggc gag tcc ttc ccg ccc 288 Pro Gln Ala Ala Gly Phe Pro Gly Ala Gly Glu Ser Phe Pro Pro 85 90 95 gcg gac gcc gag ggc tac cag ccg ggc gag ggc tac gcc gcc ccg gac 336 Ala Asp Ala Glu Gly Tyr Gln Pro Gly Glu Gly Tyr Ala Ala Pro Asp 100

									2							
													gcg Ala			384
													aac Asn			432
													atc Ile			480
													gtg Val			528
													gtc Val 190			576
													cag Gln			624
													cac His			672
													tca Ser			720
													gtg Val			768
													ctg Leu 270			816
													gct Ala			864
													gtg Val			912
													aac Asn			960
													tct Ser			1008
acc Thr	agc Ser	atc Ile	ccc Pro 340	tcc Ser	ccg Pro	cct Pro	gga Gly	ccc Pro 345	aac Asn	tgt Cys	caa Gln	ttc Phe	ctt Leu 350	ggg Gly	gga Gly	1056

_						cta Leu			_			-		_	_	1104
			-			ggc Gly 375	_		_	_		_		_	_	1152
		_		_		cgg Arg	_		_			_			_	1200
-	-	_	_	_		gca Ala		_			_					1248
						cag Gln			_	_			_			1296
						cct Pro										1344
_		_			_	ccc Pro 455	_	_								1392
					_	cag Gln				_						1440
-				_	-	tcc Ser	_	-			_		-		-	1488
						tcc Ser										1536
		-		-		tct Ser			_	_	_	_	_		_	1584
		aac Asn				aac Asn 535	tga									1608

<210> 2 <211> 535 <212> PRT

<213> Homo sapiens

<400> 2

Met Gly Ile Val Glu Pro Gly Cys Gly Asp Met Leu Thr Gly Thr Glu 1 5 10 15

Pro Met Pro Gly Ser Asp Glu Gly Arg Ala Pro Gly Ala Asp Pro Gln 20 25 30

His Arg Tyr Phe Tyr Pro Glu Pro Gly Ala Gln Asp Ala Asp Glu Arg 35 40 45

Arg Gly Gly Ser Leu Gly Ser Pro Tyr Pro Gly Gly Ala Leu Val 50 55 60

Pro Ala Pro Pro Ser Arg Phe Leu Gly Ala Tyr Ala Tyr Pro Pro Arg
65 70 75 80

Pro Gln Ala Ala Gly Phe Pro Gly Ala Gly Glu Ser Phe Pro Pro Pro 85 90 95

Ala Asp Ala Glu Gly Tyr Gln Pro Gly Glu Gly Tyr Ala Ala Pro Asp 100 105 110

Pro Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro 115 120 125

Ala Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Asn Asn His 130 135 140

Leu Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr 145 150 155 160

Lys Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly
165 170 175

Leu Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val 180 185 190

Asp Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly 195 200 205

Lys Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp 210 215 220

Ser Pro Asn Thr Gly Ala His Trp Met Arg Gln Glu Val Ser Phe Gly 225 230 235 240

Lys Leu Lys Leu Thr Asn Asn Lys Gly Ala Ser Asn Asn Val Thr Gln 245 250 255

Met Ile Val Leu Gln Ser Leu His Lys Tyr Gln Pro Arg Leu His Ile 260 265 270

Val Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Asn Ala Ser Asn 275 280 285

Thr His Ile Phe Thr Phe Gln Glu Thr Gln Phe Ile Ala Val Thr Ala 290 295 300

Tyr Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe 305 310 315 320

Ala Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Thr Ser Val Asp 325 330 335

Thr Ser Ile Pro Ser Pro Pro Gly Pro Asn Cys Gln Phe Leu Gly Gly 340 345 350

Asp His Tyr Ser Pro Leu Leu Pro Asn Gln Tyr Pro Val Pro Ser Arg 360 Phe Tyr Pro Asp Leu Pro Gly Gln Ala Lys Asp Val Val Pro Gln Ala Tyr Trp Leu Gly Ala Pro Arg Asp His Ser Tyr Glu Ala Glu Phe Arg 395 Ala Val Ser Met Lys Pro Ala Phe Leu Pro Ser Ala Pro Gly Pro Thr 405 Met Ser Tyr Tyr Arg Gly Gln Glu Val Leu Ala Pro Gly Ala Gly Trp 420 425 Pro Val Ala Pro Gln Tyr Pro Pro Lys Met Gly Pro Ala Ser Trp Phe 440 Arg Pro Met Arg Thr Leu Pro Met Glu Pro Gly Pro Gly Ser Glu 450 455 Gly Arg Gly Pro Glu Asp Gln Gly Pro Pro Leu Val Trp Thr Glu Ile 470 Ala Pro Ile Arg Pro Glu Ser Ser Asp Ser Gly Leu Gly Glu Gly Asp 485 490 Ser Lys Arg Arg Val Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser 500 505 Ser Pro Ala Gly Ala Pro Ser Pro Phe Asp Lys Glu Ala Glu Gly Gln 520 Phe Tyr Asn Tyr Phe Pro Asn 530 <210> 3 <211> 1593 <212> DNA <213> Mus musculus <220> <221> CDS <222> (1)..(1590) <400> 3 atg ggc atc gtg gag ccg ggc tgc gga gac atg ctg acc ggc acc gag 48 Met Gly Ile Val Glu Pro Gly Cys Gly Asp Met Leu Thr Gly Thr Glu ccg atg ccg agt gac gag ggc cgg ggg ccc gga gcg gac caa cag cat 96 Pro Met Pro Ser Asp Glu Gly Arg Gly Pro Gly Ala Asp Gln Gln His 20

cgt ttc ttc tat ccc gag ccg ggc gca cag gac ccg acc gat cgc cgc Arg Phe Phe Tyr Pro Glu Pro Gly Ala Gln Asp Pro Thr Asp Arg Arg

								·						
						ccc Pro								192
_		_		_		gga Gly			_		_		 _	240
_		_				cct Pro		_			_	_		288
						gtg Val								336
_						ccg Pro 120	_		-		_	_		384
	_				 _	ctg Leu	_	_			_		_	432
_			_		_	cac His	_		-	_			_	480
						ttc Phe								528
						atg Met								576
_					_	agc Ser 200		_			_	_	 _	624
						aac Asn								672
				_		atg Met	_	_	_	_			 -	720
						Gly ggg								768
	-	_	_			aag Lys		_			-			816
						gag Glu 280								864

								•						
					gag Glu 295									912
_		-			cag Gln	_			_				-	960
					ttt Phe			_		_	_	_	_	1008
_	_		_		gga Gly			_		_			-	1056
					tcc Ser		_			_	_	_		1104
		_			cag Gln 375		_	_	_		_			1152
					gaa Glu									1200
					ctc Leu									1248
					gac Asp									1296
					ccc Pro									1344
					atg Met 455									1392
					ctg Leu									1440
					cta Leu									1488
					agt Ser									1536
					gaa Glu									1584

ccc aac tga

1593

Pro Asn 530 <210> 4 <211> 530 <212> PRT <213> Mus musculus <400> 4 Met Gly Ile Val Glu Pro Gly Cys Gly Asp Met Leu Thr Gly Thr Glu Pro Met Pro Ser Asp Glu Gly Arg Gly Pro Gly Ala Asp Gln Gln His Arg Phe Phe Tyr Pro Glu Pro Gly Ala Gln Asp Pro Thr Asp Arg Arg Ala Gly Ser Ser Leu Gly Thr Pro Tyr Ser Gly Gly Ala Leu Val Pro Ala Ala Pro Gly Arg Phe Leu Gly Ser Phe Ala Tyr Pro Pro Arg Ala Gln Val Ala Gly Phe Pro Gly Pro Gly Glu Phe Phe Pro Pro Ala Gly Ala Glu Gly Tyr Pro Pro Val Asp Gly Tyr Pro Ala Pro Asp Pro Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro Ala Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Ser Asn His Leu Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr Lys 155 Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly Leu 170 Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val Asp 185 Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly Lys Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp Ser Pro Asn Thr Gly Ala His Trp Met Arg Gln Glu Val Ser Phe Gly Lys Leu Lys Leu Thr Asn Asn Lys Gly Ala Ser Asn Asn Val Thr Gln Met 250 Ile Val Leu Gln Ser Leu His Lys Tyr Gln Pro Arg Leu His Ile Val 265

Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Ser Ala Ser Asn Thr His Val Phe Thr Phe Gln Glu Thr Gln Phe Ile Ala Val Thr Ala Tyr Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe Ala 310 315 Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Ala Ser Val Asp Thr 325 330 335 Ser Val Pro Ser Pro Pro Gly Pro Asn Cys Gln Leu Leu Gly Gly Asp 340 345 Pro Phe Ser Pro Leu Leu Ser Asn Gln Tyr Pro Val Pro Ser Arg Phe 360 Tyr Pro Asp Leu Pro Gly Gln Pro Lys Asp Met Ile Ser Gln Pro Tyr 375 Trp Leu Gly Thr Pro Arg Glu His Ser Tyr Glu Ala Glu Phe Arg Ala 385 390 395 Val Ser Met Lys Pro Thr Leu Leu Pro Ser Ala Pro Gly Pro Thr Val Pro Tyr Tyr Arg Gly Gln Asp Val Leu Ala Pro Gly Ala Gly Trp Pro Val Ala Pro Gln Tyr Pro Pro Lys Met Ser Pro Ala Gly Trp Phe Arg Pro Met Arg Thr Leu Pro Met Asp Pro Gly Leu Gly Ser Ser Glu Glu Gln Gly Ser Ser Pro Ser Leu Trp Pro Glu Val Thr Ser Leu Gln Pro Glu Pro Ser Asp Ser Gly Leu Gly Glu Gly Asp Thr Lys Arg Arg Arg Ile Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser Pro Ala Gly Ala Pro Ser Pro Phe Asp Lys Glu Thr Glu Gly Gln Phe Tyr Asn Tyr Phe 520 Pro Asn 530

<210> 5 <211> 27

<212> DNA

<213> Artificial Sequence

<400> 5 gggaatttca cacctaggtg aaattcc

<210> 6 <211> 24 <212> DNA <213> Artificial Sequence	
<400> 6 aatttcacac ctaggtgtga aatt	24
<210> 7 <211> 24 <212> DNA <213> Artificial Sequence	
<400> 7 gagctatcac ctaagtgtgg gcta	24
<210> 8 <211> 24 <212> DNA <213> Artificial Sequence	
<400> 8 aaactgccac ctaagtgtgg gcta	24
<210> 9 <211> 24 <212> DNA <213> Artificial Sequence	
<400> 9	24